



PCT

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/937,100A

DATE: 01/29/2003

TIME: 15:12:54

P.6

Input Set : A:\Merk2309.app

Output Set: N:\CRF4\01292003\I937100A.raw

3 <110> APPLICANT: THIERAUCH, KARL-HEINZ
 4 GLIENKE, JENS
 5 HINZMAN, BERND
 6 PILARSKY, CHRISTIAN
 8 <120> TITLE OF INVENTION: PROTEIN ISOLATION AND ANALYSIS
 10 <130> FILE REFERENCE: MERCK 2309
 12 <140> CURRENT APPLICATION NUMBER: 09/937,100A
 C--> 13 <141> CURRENT FILING DATE: 2002-04-22
 15 <150> PRIOR APPLICATION NUMBER: PCT/GB00/01015
 16 <151> PRIOR FILING DATE: 2000-03-17
 18 <150> PRIOR APPLICATION NUMBER: 9906551.8 GB
 19 <151> PRIOR FILING DATE: 1999-03-23
 21 <150> PRIOR APPLICATION NUMBER: 9907057.5 GB
 22 <151> PRIOR FILING DATE: 1999-03-29
 24 <150> PRIOR APPLICATION NUMBER: 9907641.6 GB
 25 <151> PRIOR FILING DATE: 1999-04-06
 27 <150> PRIOR APPLICATION NUMBER: 9914874.4 GB
 28 <151> PRIOR FILING DATE: 1999-06-28
 30 <150> PRIOR APPLICATION NUMBER: 9915363.7 GB
 31 <151> PRIOR FILING DATE: 1999-07-02
 33 <150> PRIOR APPLICATION NUMBER: 9915677.0 GB
 34 <151> PRIOR FILING DATE: 1999-07-06
 36 <150> PRIOR APPLICATION NUMBER: 9916511.0 GB
 37 <151> PRIOR FILING DATE: 1999-07-14
 39 <150> PRIOR APPLICATION NUMBER: 9920503.1 GB
 40 <151> PRIOR FILING DATE: 1999-08-31
 42 <150> PRIOR APPLICATION NUMBER: 9922285.3 GB
 43 <151> PRIOR FILING DATE: 1999-09-21
 45 <160> NUMBER OF SEQ ID NOS: 69
 47 <170> SOFTWARE: PatentIn Ver. 2.1
 49 <210> SEQ ID NO: 1
 50 <211> LENGTH: 24
 51 <212> TYPE: DNA
 52 <213> ORGANISM: Artificial Sequence
 54 <220> FEATURE:
 55 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
 56 oligonucleotide for an 8 amino acid barcode peptide
 58 <220> FEATURE:
 59 <221> NAME/KEY: CDS
 60 <222> LOCATION: (1)..(24)
 61 <223> OTHER INFORMATION:
 63 <220> FEATURE:
 64 <221> NAME/KEY: misc_feature

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65 <222> LOCATION: (1)..(1)
66 <223> OTHER INFORMATION: n=a,t,g,c
W--> 68 <220>
69 <221> NAME/KEY: misc_feature
70 <222> LOCATION: (4)..(4)
71 <223> OTHER INFORMATION: n=a,t,g,c
73 <220> FEATURE:
74 <221> NAME/KEY: misc_feature
75 <222> LOCATION: (7)..(7)
76 <223> OTHER INFORMATION: n=a,t,g,c
78 <220> FEATURE:
79 <221> NAME/KEY: misc_feature
80 <222> LOCATION: (10)..(10)
81 <223> OTHER INFORMATION: n=a,t,g,c
83 <220> FEATURE:
84 <221> NAME/KEY: misc_feature
85 <222> LOCATION: (14)..(14)
86 <223> OTHER INFORMATION: k=g,t
88 <220> FEATURE:
89 <221> NAME/KEY: misc_feature
90 <222> LOCATION: (16)..(16)
91 <223> OTHER INFORMATION: v=a,g,c
93 <220> FEATURE:
94 <221> NAME/KEY: misc_feature
95 <222> LOCATION: (20)..(20)
96 <223> OTHER INFORMATION: n=a,g,t,c
98 <220> FEATURE:
99 <221> NAME/KEY: misc_feature
100 <222> LOCATION: (21)..(21)
101 <223> OTHER INFORMATION: v=a,g,c
103 <220> FEATURE:
104 <221> NAME/KEY: misc_feature
105 <222> LOCATION: (23)..(23)
106 <223> OTHER INFORMATION: n=a,t,g,c
108 <400> SEQUENCE: 1
W--> 109 nac ncc ngg ntg tkc vag gnv cnt 24
W--> 110 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
111 1 5
114 <210> SEQ ID NO: 2
115 <211> LENGTH: 8
116 <212> TYPE: PRT
117 <213> ORGANISM: Artificial Sequence
119 <220> FEATURE:
120 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
121 barcode peptide
123 <220> FEATURE:
124 <221> NAME/KEY: misc_feature
125 <222> LOCATION: (1)..(1)
126 <223> OTHER INFORMATION: The 'Xaa' at location 1 stands for Asn, Asp, His, or Tyr.

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128 <220> FEATURE:
129 <221> NAME/KEY: misc_feature
130 <222> LOCATION: (2)..(2)
131 <223> OTHER INFORMATION: The 'Xaa' at location 2 stands for Thr, Ala, Pro, or Ser.
133 <220> FEATURE:
134 <221> NAME/KEY: misc_feature
135 <222> LOCATION: (3)..(3)
136 <223> OTHER INFORMATION: The 'Xaa' at location 3 stands for Arg, Gly, or Trp.
138 <220> FEATURE:
139 <221> NAME/KEY: misc_feature
140 <222> LOCATION: (4)..(4)
141 <223> OTHER INFORMATION: The 'Xaa' at location 4 stands for Met, Val, or Leu.
143 <220> FEATURE:
144 <221> NAME/KEY: misc_feature
145 <222> LOCATION: (5)..(5)
146 <223> OTHER INFORMATION: The 'Xaa' at location 5 stands for Cys, or Phe.
148 <220> FEATURE:
149 <221> NAME/KEY: misc_feature
150 <222> LOCATION: (6)..(6)
151 <223> OTHER INFORMATION: The 'Xaa' at location 6 stands for Lys, Glu, or Gln.
153 <220> FEATURE:
154 <221> NAME/KEY: misc_feature
155 <222> LOCATION: (7)..(7)
156 <223> OTHER INFORMATION: The 'Xaa' at location 7 stands for Glu, Asp, Gly, Ala, or
Val.
158 <220> FEATURE:
159 <221> NAME/KEY: misc_feature
160 <222> LOCATION: (8)..(8)
161 <223> OTHER INFORMATION: The 'Xaa' at location 8 stands for His, Arg, Pro, or Leu.
163 <400> SEQUENCE: 2
W--> 164 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
165 1 5
168 <210> SEQ ID NO: 3
169 <211> LENGTH: 14
170 <212> TYPE: PRT
171 <213> ORGANISM: Artificial Sequence
173 <220> FEATURE:
174 <223> OTHER INFORMATION: Description of Artificial Sequence: Linker
175 peptide
177 <400> SEQUENCE: 3
178 Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Val Asp
179 1 5 10
182 <210> SEQ ID NO: 4
183 <211> LENGTH: 8
184 <212> TYPE: PRT
185 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Description of Artificial Sequence: Flag
189 epitope peptide
191 <400> SEQUENCE: 4

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192 Met Asp Tyr Lys Asp Asp Asp Lys
193 1 5
196 <210> SEQ ID NO: 5
197 <211> LENGTH: 53
198 <212> TYPE: DNA
199 <213> ORGANISM: Artificial Sequence
201 <220> FEATURE:
202 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
203 RD5' Flag
205 <400> SEQUENCE: 5
206 gcg gatccca tatg gactac aaagacgatg acgacaaaca ggtgcagctg cag 53
209 <210> SEQ ID NO: 6
210 <211> LENGTH: 35
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
216 RD3'
218 <400> SEQUENCE: 6
219 gcgaattcgt ggtggtggtg gtggtgtgac tctcc 35
222 <210> SEQ ID NO: 7
223 <211> LENGTH: 50
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
229 Foslfor
231 <400> SEQUENCE: 7
232 atggaattcc tcgagaccga caccctacag gcggaaaccg accagctgga 50
235 <210> SEQ ID NO: 8
236 <211> LENGTH: 50
237 <212> TYPE: DNA
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
242 Fos80rev
244 <400> SEQUENCE: 8
245 tcgcgatttc ggtttgcagc gcggattttt cgtcttccag ctggtcggtt 50
248 <210> SEQ ID NO: 9
249 <211> LENGTH: 50
250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
255 Fos71for
257 <400> SEQUENCE: 9
258 aaaccgaaat cgcaacctg ctgaaagaaa aagaaaagct ggagttcatc 50
261 <210> SEQ ID NO: 10
262 <211> LENGTH: 50

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263 <212> TYPE: DNA
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
268     Fos155rev
270 <400> SEQUENCE: 10
271 ggaagcttga attccgccgg acggtgtgcc gccaggatga actccagctt      50
274 <210> SEQ ID NO: 11
275 <211> LENGTH: 18
276 <212> TYPE: DNA
277 <213> ORGANISM: Artificial Sequence
279 <220> FEATURE:
280 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
281     Fos1 fS
283 <400> SEQUENCE: 11
284 atggaattcc tcgagacc      18
287 <210> SEQ ID NO: 12
288 <211> LENGTH: 18
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
294     Fos155 rS
296 <400> SEQUENCE: 12
297 ggaagcttga attccgcc      18
300 <210> SEQ ID NO: 13
301 <211> LENGTH: 28
302 <212> TYPE: DNA
303 <213> ORGANISM: Artificial Sequence
305 <220> FEATURE:
306 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer I
307     for 340 VH amplification
309 <400> SEQUENCE: 13
310 cagctgcagg agtctggggg aggcttag      28
313 <210> SEQ ID NO: 14
314 <211> LENGTH: 36
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial Sequence
318 <220> FEATURE:
319 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer II
320     for 340 VH amplification
322 <400> SEQUENCE: 14
323 tcagtagacg gtgaccgagg ttccttgacc ccagta      36
326 <210> SEQ ID NO: 15
327 <211> LENGTH: 26
328 <212> TYPE: DNA
329 <213> ORGANISM: Artificial Sequence
331 <220> FEATURE:
332 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer I

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RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1,4,7,10,20,23
Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:2; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:27; N Pos. 21,24
Seq#:28; N Pos. 21,24
Seq#:29; N Pos. 21,24
Seq#:30; N Pos. 21,24
Seq#:31; N Pos. 21,24
Seq#:32; N Pos. 7,10
Seq#:33; N Pos. 7,10
Seq#:34; N Pos. 7,10
Seq#:35; N Pos. 7,10
Seq#:36; N Pos. 7,10
Seq#:58; N Pos. 31,32,34,35,37,38,40,41,43,44,46,47,49,50
Seq#:58; Xaa Pos. 11,12,13,14,15,16,17
Seq#:59; Xaa Pos. 11,12,13,14,15,16,17
Seq#:60; N Pos. 28,29,31,32,34,35,37,38,40,41,43,44,46,47
Seq#:61; N Pos. 14,15,17,18,20,21,23,24,26,27,29,30,32,33,35,36,38,39,41,42
Seq#:61; Xaa Pos. 5,6,7,8,9,10,11,12,13,14
Seq#:62; Xaa Pos. 5,6,7,8,9,10,11,12,13,14
Seq#:63; N Pos. 29,30,32,33,35,36,38,39,41,42,44,45,47,48,50,51,53,54,56,57
Seq#:64; N Pos. 15,18,19,22,25,28,38,41
Seq#:64; Xaa Pos. 7,8,9,10,11,12,13,14
Seq#:65; Xaa Pos. 7,8,9,10,11,12,13,14
Seq#:66; N Pos. 14,17,27,30,33,36
Seq#:67; N Pos. 15,18,19,22,25,28,38,41,48,51,52,55,58,61,71,74
Seq#:67; Xaa Pos. 7,8,9,10,11,12,13,14,18,19,20,21,22,23,24,25
Seq#:68; Xaa Pos. 7,8,9,10,11,12,13,14,18,19,20,21,22,23,24,25
Seq#:69; N Pos. 14,17,27,30,33,36,47,50,60,63,66,69

VERIFICATION SUMMARY

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Input Set : A:\Merk2309.app

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:68 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:61
L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:24
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:640 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:709 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:1017 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:58,Line#:1010
L:1083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
L:1084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:48
L:1087 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:48
L:1088 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:77
L:1151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:16
L:1239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0
L:1263 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:61,Line#:1256
L:1359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
L:1360 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:49
L:1448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0
L:1566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0
L:1589 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:64,Line#:1582
L:1645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
L:1646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:48
L:1703 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
L:1762 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
L:1784 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:67,Line#:1777
L:1890 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:0
L:1891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:48
L:1894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:48
L:1895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:87
L:1989 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:0
L:1992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:16
L:2098 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:0
L:2099 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:60